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CAGGTCCAAC	TGCAGGAGTC	AGGGGGAGAT	CTAGTGCAGC	CTGGAAGGTC	50
GTCCAGGTTG	ACGTCCTCAG	TCCCCCTCTA	GATCACGTCG	GACCTTCCAG	
Q V Q L	Q E S	G G D	L V Q P	G R S	
TCTGAAACTT	TCCTGTGTAG	CCTCTGGATT	CACATTCAGT	AATTACTGGA	100
AGACTTTGAA	AGGACACATC	GGAGACCTAA	GTGTAAGTCA	TTAATGACCT	
L K L	S C V A	S G F	T F S	<u>N Y W M</u>	
CDR1					
TGACTTGAT	CCGCCAGGCT	CCAGGGGAGG	GTCTTGAATG	GGTTGCGTCC	150
ACTGAACCTA	GGCGGTCCGA	GGTCCCCCTCC	CAGAACTTAC	CCAACGCAGG	
<u>T</u> W I	R Q A	P G E G	L E W	V A <u>S</u>	
ATTACTAGTA	CTGGTGGTGG	GACTTACCAT	GCAGAGTCTG	TGAAGGGCCG	200
TAATGATCAT	GACCACCACC	CTGAATGGTA	CGTCTCAGAC	ACTTCCCGGC	
<u>I</u> T S T	G G G	<u>T Y H</u>	A E S V	K G R	
CDR2					
ATTCACTATC	TCCAGAGATA	ATTCAAAAAG	CACCCTGTAC	CTGCAAATGA	250
TAAGTGATAG	AGGTCTCTAT	TAAGTTTTTC	GTGGGACATG	GACGTTTACT	
F T I	S R D N	S K S	T L Y	L Q M N	
ACAGTCTGAG	GCCTGAGGAC	ACGGCCACTT	ATTACTGTTC	AAGAGATGAC	300
TGTCAGACTC	CGGACTCCTG	TGCCGGTGAA	TAATGACAAG	TTCTCTACTG	
S L R	P E D	T A T Y	Y C S	R <u>D D</u>	
TACGGAGGAC	AGAGCACCTA	TGTTATGGAT	GCCTGGGGTC	AGGGATCTTC	350
ATGCCTCCTG	TCTCGTGGAT	ACAATACCTA	CGGACCCAG	TCCCTAGAAG	
<u>Y G G Q</u>	<u>S T Y</u>	<u>V M D A</u>	W G Q	G S S	
CDR3					
GGTCAACGTC	TCCTCA				366
CCAGTGGCAG	AGGAGT				
V T V	S S				

Fig. 7